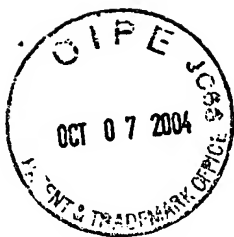




		Section 1					
		(1) 1	10	20	30	40	52
ApoB 1-269	(1)	-----MDPPRPALLALLA[P]LLLLLLAGAR[EE][LE					
CD1d human	(1)	MGCLLFLLLWALLQAWGSAEVPQRLFPLRCLQ[S]FANSSWTRTD[LAW][GE					
		Section 2					
		(53) 53	60	70	80	90	104
ApoB 1-269	(34)	NV[LVC	PKD[TRF[HL[K[YN-----YEA	E--S--SSG	VP	GTA[SRS	
CD1d human	(53)	LQ[HSW	SND[DTV[SL[P[QGT	FSDQ	QWETLQHIF	RVRSS	FTRDVK[FAK
		Section 3					
		(105) 105	110	120	130	140	156
ApoB 1-269	(73)	ATR[NCK[ELEV	PQLCSFILK--[S-----QCT	LKE	VYGFNP		
CD1d human	(105)	MLR[SY	P[ELQVS	AGCEVHPGNA[NNFF	HVA	FQ	GKDILSFQGTSWEPTQEAP
		Section 4					
		(157) 157	170	180	190	208	
ApoB 1-269	(108)	EGKALLK	TKNS[E[AAAM	SR[ELKL	AIPE	GKQVFLYPE	KDEPTYILN
CD1d human	(157)	LWVNL	AIQVLNQ[K[TRE	TVQ[LLNG	TC	PQ	FVSGLL
							ESGKSE-----LKKQV
		Section 5					
		(209) 209	220	230	240	250	260
ApoB 1-269	(160)	IISALL	VPPE[E	EAKQ[LF[DTV	YGNCSTHFT	VKTR[GN-----[A-	
CD1d human	(204)	KPKAW	LSRGP[P	GPGR[LL[CHV	SGFY	PKPVV	WKWM[GEQE
							QQGTQPGD[LP
		Section 6					
		(261) 261	270	280	290	300	312
ApoB 1-269	(201)	-----T-----[STER	DLGQCD	RFKPI	RTGIS	P[ALIK	GMRPL
CD1d human	(256)	NADET	WYLRATL[VAGE	AAGL	SCR	VKHSS	LEGQD[VLY
							WGG[YTSMGLI[L
		Section 7					
		(313) 313	320	330	343		
ApoB 1-269	(242)	[QS-----CQ[TL	DAKR	KHVAE[CKEQ	SEQ ID NO: 11		
CD1d human	(308)	[VLAC	LLFLLIVG[TSR	FRQ	TSYQ[L---	SEQ ID NO: 12	

Fig. 12



		Section 1					
		(1) 1	10	20	30	40	52
ApoB 512-721	(1)	-----K					
CD1d human	(1)	MGCLLFLLLWALLQAWGSAEVPQRLFPLRCLQISSFANSSWTRTDGLAWLGE					
		Section 2					
		(53) 53	60	70	80	90	104
ApoB 512-721	(2)	CVQS	KPSLMIQKA	QALRKMEPKDKDQEVLLQTF	LDDASPGD	R	AA
CD1d human	(53)	LQTH	WSNDSDTVR	KPWSQGTFS	DQWETLQHI	FRVYRSSFT	D
		Section 3					
		(105) 105	110	120	130	140	156
ApoB 512-721	(54)	RS--PSQAD	NKIVQ	LPWEQNE	-----QVKN	VA	HIAN
CD1d human	(105)	RLSYPLELQ	SAGCE	HPGNASNNFFH	VAFQGKDILS	FQGT	S
		Section 4					
		(157) 157	170	180	190	208	
ApoB 512-721	(89)	LNSEE	D	QDLKKLV	EA	KESQLPTVMDFRKFSRN	-----YQLYKSV
CD1d human	(157)	WVNLA	Q	LNQDKWT	ET	QWLLNGT	CPQFVSGLL
		Section 5					
		(209) 209	220	230	240	250	260
ApoB 512-721	(135)	PSLDP	S	KIEGNL	FDPNN	LPK--ESM	TTLTAFGFAS
CD1d human	(209)	LSRGP	P	PGRLLL	CHVSG	YPKPVWVKW	GEQEQQGTQP
		Section 6					
		(261) 261	270	280	290	300	312
ApoB 512-721	(184)	KGFEPTL	A	FGKQ	-----FFP	SVNKALYW	-----
CD1d human	(261)	WYLRATL	V	AGEA	GLSCRVKHSSL	GQDIVLYW	GGSYTSMGLIALAVLAC
		Section 7					
		(313) 313	320	335			
ApoB 512-721	(211)	-----					
CD1d human	(313)	LLFLLIVGFTSRFKRQTSYQGV					
					SEQ ID NO: 13		
					SEQ ID NO: 15		

Fig. 13



		Section 1						
		(1)	1	10	20	30	40	52
ApoB 270-570	(1)	-----FSYNNKYGMVAQVTQTLKLEDTPKINSRFFGEGT						KMGLA
CD1d human	(1)	MGCLLFLLLWALLQAWGSAEVPQRLFPLRCLQISSFANSSWT						TDGLA
		Section 2						
		(53)	53	60	70	80	90	104
ApoB 270-570	(45)	KSTSPPK	AEAV		LQELKKLTIS	Q	IQRA	LFN
CD1d human	(53)	LQTHSWS	DSDT		LKPWSQGTFS	Q	WETL	HIF
		Section 3						
		(105)	105	110	120	130	140	156
ApoB 270-570	(97)	LPQLIE	SSP		LQALVQCG	PQCSTHI	Q	LK
CD1d human	(105)	LRLSYP	ELQ		AGCEVHPG	ASNNFFH	A	QG
		Section 4						
		(157)	157	170	180	190	208	
ApoB 270-570	(149)	PEPSAQQ	REIFNMAR		QSRATLYALSHAVNNYHK	NP	GTQEL	DIAN
CD1d human	(157)	WVNLAIQ	LNQDKWTR		TVQWLLNGT	CPQFVSGLLE	GK	ELKKQ
		Section 5						
		(209)	209	220	230	240	250	260
ApoB 270-570	(201)	LMEQIQDD	-----C		GDEDYTYL	LR	IGNMGQTMEQLTPELKS	ILK
CD1d human	(209)	LSRGPSPGPGRLLL	VCHV		GFYPKPVW	KW	RGEQEQQGTQPGDILPN	DET
		Section 6						
		(261)	261	270	280	290	300	312
ApoB 270-570	(244)	CVQSTKPS		IQKAAIQ	LRKMEPKDKDQ		LQTFLDDASPGDKRLAAYLM	
CD1d human	(261)	WYLRATLD		AGEAAGL	CRVKHSSLEGQ		LYWGGSYTSMGLIALAVLAC	
		Section 7						
		(313)	313	320	335			
ApoB 270-570	(296)	L	RSPS	-----	SEQ ID NO: 14			
CD1d human	(313)	L	FLLI	VGFTSRFKRQTSYQGV	SEQ ID NO: 16			

Fig. 14